

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readables form:

Application Serial Number: 09/707,738A

Source: /600

Date Processed by STIC: 8/14/2003

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRE SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212:

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry directly to:
 - U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

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DATE: 08/14/2003

1600

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PATENT APPLICATION: US/09/707,738A TIME: 15:30:47
                      Input Set : A:\-62-5.app
                      Output Set: N:\CRF4\08142003\1707738A.raw
       4 <110> APPLICANT: Sette, Alessandro
               Gaeta, Federico
               Grey, Howard M.
               Sidney, John
              Alexander, Jeffery L.
               Epimmune Inc.
     11 <120> TITLE OF INVENTION: Induction of Immune Response Against
              Desired Determinants
     14 <130> FILE REFERENCE: 018623-006250US
     16 <140> CURRENT APPLICATION NUMBER: US 09/707,738A
     17 <141> CURRENT FILING DATE: 2000-11-06
     19 <150> PRIOR APPLICATION NUMBER: US 08/121,101
     20 <151> PRIOR FILING DATE: 1993-09-14
     22 <150> PRIOR APPLICATION NUMBER: US 08/305,871
     23 <151> PRIOR FILING DATE: 1994-09-14
     25 <150> PRIOR APPLICATION NUMBER: US 08/485,218
     26 <151> PRIOR FILING DATE: 1995-06-07
     28 <150> PRIOR APPLICATION NUMBER: US 60/010,510
     29 <151> PRIOR FILING DATE: 1996-01-24
     31 <150> PRIOR APPLICATION NUMBER: US 08/788,822
     32 <151> PRIOR FILING DATE: 1997-01-23
     34 <150> PRIOR APPLICATION NUMBER: US 09/310,462
                                                                  Does Not Comply
     35 <151> PRIOR FILING DATE: 1999-05-12
E--> 37 <160> NUMBER OF SEQ ID NOS: (25)27 (\rho,3)
                                                              Corrected Diskette Needed
                                                              pp1-3
     39 <170> SOFTWARE: FastSEQ for Windows Version 3.0
     453 <210> SEQ ID NO: 26
454 <211> LENGTH: (13) 16 Shown (\rho. 2)
     455 <212> TYPE: PRT
     456 <213> ORGANISM: Artificial Sequence
     458 <220> FEATURE:
     459 <223> OTHER INFORMATION: peptide binds more than one DR allele
W--> 461 <221> NAME/KEY: MOD_RES
     462 <222> LOCATION: (1)...(1)
     463 <223> OTHER INFORMATION: Xaa = any D- or L-amino acid
W--> 465 <221> MOD_RES
     466 <222> LOCATION: (2)...(2)
     467 <223> OTHER INFORMATION: Xaa = Ala or Lys
W--> 469 <221> MOD_RES
     470 <222> LOCATION: (3)...(3)
```

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING DATE: 08/14/2003 PATENT APPLICATION: US/09/707,738A TIME: 15:30:48

Input Set : A:\-62-5.app

Output Set: N:\CRF4\08142003\1707738A.raw

471 <223> OTHER INFORMATION: Xaa = cyclohexylalanine, Tyr or Phe W--> 473 <221> MOD_RES 474 <222> LOCATION: (4)...(6) 475 <223> OTHER INFORMATION: Xaa = Ala, Ile, Ser or Val W--> 477 <221> MOD_RES 478 <222> LOCATION: (7)...(7) 479 <223> OTHER INFORMATION: Xaa = Ala, Ile, Ser or Val, Xaa at position 7 may be present or absent 480 W--> 482 <221> MOD_RES 483 <222> LOCATION: (12)...(13) 484 <223> OTHER INFORMATION: Xaa = Ala, Ser or Val W--> 486 <221> MOD_RES 487 <222> LOCATION: (14)...(15) 488 <223> OTHER INFORMATION: Xaa = Ala, Ser or Val, Xaa at positions 14 and 15 489 may be present or absent W--> 491 <221> MOD_RES 492 <222> LOCATION: (16)...(16) 493 <223> OTHER INFORMATION: Xaa = any D- or L-amino acid W--> 495 <400> 26 W--> 496 Xaa Xaa Xaa Xaa Xaa Xaa Trp Thr Leu Lys Xaa Xaa Xaa Xaa Xaa 5 E--> 497 1 15

see p3

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<210> 27 last sequence in submitted file
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> pan DR binding peptide binding core
<400> 27
Trp Thr Leu Lys

Use of n and/or Xaa has been detected in the Sequence Listing.
Review the S requence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xss.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/707,738A

DATE: 08/14/2003
TIME: 15:30:49

Input Set : A:\-62-5.app

Output Set: N:\CRF4\08142003\1707738A.raw

```
L:329 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:333 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:337 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:350 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:354 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:358 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:362 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:366 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:370 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:374 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:387 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:391 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:395 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:399 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:403 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:407 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:411 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:424 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:428 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:432 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:436 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:440 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:444 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:448 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:449 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:461 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:465 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:469 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:473 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:477 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:482 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:486 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:491 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:495 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:497 M:252 E: No. of Seq. differs, <211> LENGTH:Input:13 Found:16 SEQ:26
L:37 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (25) Counted (27)
```